SEQUENCE LISTING

<110> Katherine Galvin and Laura A. Rudolph-Owen

<120> METHODS AND COMPOSITIONS FOR THE DIAGNOSIS
      AND TREATMENT OF CARDIOVASCULAR AND TUMORIGENIC
      DISEASE USING 4941

<130> MNI-094

<140> 09/635,521
<141> 2000-08-09

<150> 60/199,908
<151> 2000-04-26

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<170> PatentIn Ver. 2.0

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<213> Homo sapiens

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Met Ala Ser Pro Ser Leu Pro Gly Ser Asp Cys Ser Gln Ile Ile Asp
 1               5                   10                  15

cac agt cat gtc ccc gag ttt gag gtg gcc acc tgg atc aaa atc acc    96
His Ser His Val Pro Glu Phe Glu Val Ala Thr Trp Ile Lys Ile Thr
            20                  25                  30

ctt att ctg gtg tac ctg atc atc ttc gtg atg ggc ctt ctg ggg aac   144
Leu Ile Leu Val Tyr Leu Ile Ile Phe Val Met Gly Leu Leu Gly Asn
        35                  40                  45

agc gcc acc att cgg gtc acc cag gtg ctg cag aag aaa gga tac ttg   192
Ser Ala Thr Ile Arg Val Thr Gln Val Leu Gln Lys Lys Gly Tyr Leu
    50                  55                  60

cag aag gag gtg aca gac cac atg gtg agt ttg gct tgc tcg gac atc   240
Gln Lys Glu Val Thr Asp His Met Val Ser Leu Ala Cys Ser Asp Ile
65                  70                  75                  80

ttg gtg ttc ctc atc ggc atg ccc atg gag ttc tac agc atc atc tgg   288
Leu Val Phe Leu Ile Gly Met Pro Met Glu Phe Tyr Ser Ile Ile Trp
                85                  90                  95

aat ccc ctg acc acg tcc agc tac acc ctg tcc tgc aag ctg cac act   336
Asn Pro Leu Thr Thr Ser Ser Tyr Thr Leu Ser Cys Lys Leu His Thr
                100                 105                 110
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ttc ctc ttc gag gcc tgc agc tac gct acg ctg ctg cac gtg ctg aca    384
Phe Leu Phe Glu Ala Cys Ser Tyr Ala Thr Leu Leu His Val Leu Thr
        115                 120                 125

ctc agc ttt gag cgc tac atc gcc atc tgt cac ccc ttc agg tac aag    432
Leu Ser Phe Glu Arg Tyr Ile Ala Ile Cys His Pro Phe Arg Tyr Lys
        130                 135                 140

gct gtg tcg gga cct tgc cag gtg aag ctg ctg att ggc ttc gtc tgg    480
Ala Val Ser Gly Pro Cys Gln Val Lys Leu Leu Ile Gly Phe Val Trp
145                 150                 155                 160

gtc acc tcc gcc ctg gtg gca ctg ccc ttg ctg ttt gcc atg ggt act    528
Val Thr Ser Ala Leu Val Ala Leu Pro Leu Leu Phe Ala Met Gly Thr
                165                 170                 175

gag tac ccc ctg gtg aac gtg ccc agc cac cgg ggt ctc act tgc aac    576
Glu Tyr Pro Leu Val Asn Val Pro Ser His Arg Gly Leu Thr Cys Asn
                180                 185                 190

cgc tcc agc acc cgc cac cac gag cag ccc gag acc tcc aat atg tcc    624
Arg Ser Ser Thr Arg His His Glu Gln Pro Glu Thr Ser Asn Met Ser
        195                 200                 205

atc tgt acc aac ctc tcc agc cgc tgg acc gtg ttc cag tcc agc atc    672
Ile Cys Thr Asn Leu Ser Ser Arg Trp Thr Val Phe Gln Ser Ser Ile
    210                 215                 220

ttc ggc gcc ttc gtg gtc tac ctc gtg gtc ctg ctc tcc gta gcc ttc    720
Phe Gly Ala Phe Val Val Tyr Leu Val Val Leu Leu Ser Val Ala Phe
225                 230                 235                 240

atg tgc tgg aac atg atg cag gtg ctc atg aaa agc cag aag ggc tcg    768
Met Cys Trp Asn Met Met Gln Val Leu Met Lys Ser Gln Lys Gly Ser
                245                 250                 255

ctg gcc ggg ggc acg cgg cct ccg cag ctg agg aag tcc gag agc gaa    816
Leu Ala Gly Gly Thr Arg Pro Pro Gln Leu Arg Lys Ser Glu Ser Glu
                260                 265                 270

gag agc agg acc gcc agg agg cag acc atc atc ttc ctg agg ctg att    864
Glu Ser Arg Thr Ala Arg Arg Gln Thr Ile Ile Phe Leu Arg Leu Ile
        275                 280                 285

gtt gtg aca ttg gcc gta tgc tgg atg ccc aac cag att cgg agg atc    912
Val Val Thr Leu Ala Val Cys Trp Met Pro Asn Gln Ile Arg Arg Ile
        290                 295                 300

atg gct gcg gcc aaa ccc aag cac gac tgg acg agg tcc tac ttc cgg    960
Met Ala Ala Ala Lys Pro Lys His Asp Trp Thr Arg Ser Tyr Phe Arg
305                 310                 315                 320

gcg tac atg atc ctc ctc ccc ttc tcg gag acg ttt ttc tac ctc agc   1008
Ala Tyr Met Ile Leu Leu Pro Phe Ser Glu Thr Phe Phe Tyr Leu Ser
                325                 330                 335
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tcg gtc atc aac ccg ctc ctg tac acg gtg tcc tcg cag cag ttt cgg      1056
Ser Val Ile Asn Pro Leu Leu Tyr Thr Val Ser Ser Gln Gln Phe Arg
            340             345             350

cgg gtg ttc gtg cag gtg ctg tgc tgc cgc ctg tcg ctg cag cac gcc      1104
Arg Val Phe Val Gln Val Leu Cys Cys Arg Leu Ser Leu Gln His Ala
            355             360             365

aac cac gag aag cgc ctg cgc gta cat gcg cac tcc acc acc gac agc      1152
Asn His Glu Lys Arg Leu Arg Val His Ala His Ser Thr Thr Asp Ser
            370             375             380

gcc cgc ttt gtg cag cgc ccg ttg ctc ttc gcg tcc cgg cgc cag tcc      1200
Ala Arg Phe Val Gln Arg Pro Leu Leu Phe Ala Ser Arg Arg Gln Ser
385             390             395             400

tct gca agg aga act gag aag att ttc tta agc act ttt cag agc gag      1248
Ser Ala Arg Arg Thr Glu Lys Ile Phe Leu Ser Thr Phe Gln Ser Glu
            405             410             415

gcc gag ccc cag tct aag tcc cag tca ttg agt ctc gag tca cta gag      1296
Ala Glu Pro Gln Ser Lys Ser Gln Ser Leu Ser Leu Glu Ser Leu Glu
            420             425             430

ccc aac tca ggc gcg aaa cca gcc aat tct gct gca gag aat ggt ttt      1344
Pro Asn Ser Gly Ala Lys Pro Ala Asn Ser Ala Ala Glu Asn Gly Phe
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cag gag cat gaa gtt tga                                               1362
Gln Glu His Glu Val
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His Ser His Val Pro Glu Phe Glu Val Ala Thr Trp Ile Lys Ile Thr
             20             25                  30

Leu Ile Leu Val Tyr Leu Ile Ile Phe Val Met Gly Leu Leu Gly Asn
        35              40                  45

Ser Ala Thr Ile Arg Val Thr Gln Val Leu Gln Lys Lys Gly Tyr Leu
     50             55                  60

Gln Lys Glu Val Thr Asp His Met Val Ser Leu Ala Cys Ser Asp Ile
 65             70                  75                  80

Leu Val Phe Leu Ile Gly Met Pro Met Glu Phe Tyr Ser Ile Ile Trp
                85              90                  95
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Asn Pro Leu Thr Thr Ser Ser Tyr Thr Leu Ser Cys Lys Leu His Thr
          100   105   110

Phe Leu Phe Glu Ala Cys Ser Tyr Ala Thr Leu Leu His Val Leu Thr
          115   120   125

Leu Ser Phe Glu Arg Tyr Ile Ala Ile Cys His Pro Phe Arg Tyr Lys
          130   135   140

Ala Val Ser Gly Pro Cys Gln Val Lys Leu Leu Ile Gly Phe Val Trp
145   150   155   160

Val Thr Ser Ala Leu Val Ala Leu Pro Leu Leu Phe Ala Met Gly Thr
              165   170   175

Glu Tyr Pro Leu Val Asn Val Pro Ser His Arg Gly Leu Thr Cys Asn
          180   185   190

Arg Ser Ser Thr Arg His His Glu Gln Pro Glu Thr Ser Asn Met Ser
          195   200   205

Ile Cys Thr Asn Leu Ser Ser Arg Trp Thr Val Phe Gln Ser Ser Ile
          210   215   220

Phe Gly Ala Phe Val Val Tyr Leu Val Val Leu Leu Ser Val Ala Phe
225   230   235   240

Met Cys Trp Asn Met Met Gln Val Leu Met Lys Ser Gln Lys Gly Ser
              245   250   255

Leu Ala Gly Gly Thr Arg Pro Pro Gln Leu Arg Lys Ser Glu Ser Glu
              260   265   270

Glu Ser Arg Thr Ala Arg Arg Gln Thr Ile Ile Phe Leu Arg Leu Ile
          275   280   285

Val Val Thr Leu Ala Val Cys Trp Met Pro Asn Gln Ile Arg Arg Ile
          290   295   300

Met Ala Ala Ala Lys Pro Lys His Asp Trp Thr Arg Ser Tyr Phe Arg
305   310   315   320

Ala Tyr Met Ile Leu Leu Pro Phe Ser Glu Thr Phe Phe Tyr Leu Ser
              325   330   335

Ser Val Ile Asn Pro Leu Leu Tyr Thr Val Ser Ser Gln Gln Phe Arg
              340   345   350

Arg Val Phe Val Gln Val Leu Cys Cys Arg Leu Ser Leu Gln His Ala
          355   360   365

Asn His Glu Lys Arg Leu Arg Val His Ala His Ser Thr Thr Asp Ser
          370   375   380

Ala Arg Phe Val Gln Arg Pro Leu Leu Phe Ala Ser Arg Arg Gln Ser
385   390   395   400

Ser Ala Arg Arg Thr Glu Lys Ile Phe Leu Ser Thr Phe Gln Ser Glu
            405   410   415

Ala Glu Pro Gln Ser Lys Ser Gln Ser Leu Ser Leu Glu Ser Leu Glu
            420   425   430

Pro Asn Ser Gly Ala Lys Pro Ala Asn Ser Ala Ala Glu Asn Gly Phe
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Gln Glu His Glu Val
            450

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                                             Met Ala Ser Pro Ser
                                              1   5

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Leu Pro Gly Ser Asp Cys Ser Gln Ile Ile Asp His Ser His Val Pro
              10   15   20

gag ttt gag gtg gcc acc tgg atc aaa atc acc ctt att ctg gtg tac   152
Glu Phe Glu Val Ala Thr Trp Ile Lys Ile Thr Leu Ile Leu Val Tyr
              25   30   35

ctg atc atc ttc gtg atg ggc ctt ctg ggg aac agc gcc acc att cgg   200
Leu Ile Ile Phe Val Met Gly Leu Leu Gly Asn Ser Ala Thr Ile Arg
              40   45   50

gtc acc cag gtg ctg cag aag aaa gga tac ttg cag aag gag gtg aca   248
Val Thr Gln Val Leu Gln Lys Lys Gly Tyr Leu Gln Lys Glu Val Thr
      55   60   65

gac cac atg gtg agt ttg gct tgc tcg gac atc ttg gtg ttc ctc atc   296
Asp His Met Val Ser Leu Ala Cys Ser Asp Ile Leu Val Phe Leu Ile
 70   75   80   85

ggc atg ccc atg gag ttc tac agc atc atc tgg aat ccc ctg acc acg   344
Gly Met Pro Met Glu Phe Tyr Ser Ile Ile Trp Asn Pro Leu Thr Thr
                  90   95   100

tcc agc tac acc ctg tcc tgc aag ctg cac act ttc ctc ttc gag gcc   392
Ser Ser Tyr Thr Leu Ser Cys Lys Leu His Thr Phe Leu Phe Glu Ala

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tgc agc tac gct acg ctg ctg cac gtg ctg aca ctc agc ttt gag cgc    440
Cys Ser Tyr Ala Thr Leu Leu His Val Leu Thr Leu Ser Phe Glu Arg
        120                 125                 130

tac atc gcc atc tgt cac ccc ttc agg tac aag gct gtg tcg gga cct    488
Tyr Ile Ala Ile Cys His Pro Phe Arg Tyr Lys Ala Val Ser Gly Pro
        135                 140                 145

tgc cag gtg aag ctg ctg att ggc ttc gtc tgg gtc acc tcc gcc ctg    536
Cys Gln Val Lys Leu Leu Ile Gly Phe Val Trp Val Thr Ser Ala Leu
150                 155                 160                 165

gtg gca ctg ccc ttg ctg ttt gcc atg ggt act gag tac ccc ctg gtg    584
Val Ala Leu Pro Leu Leu Phe Ala Met Gly Thr Glu Tyr Pro Leu Val
                170                 175                 180

aac gtg ccc agc cac cgg ggt ctc act tgc aac cgc tcc agc acc cgc    632
Asn Val Pro Ser His Arg Gly Leu Thr Cys Asn Arg Ser Ser Thr Arg
                185                 190                 195

cac cac gag cag ccc gag acc tcc aat atg tcc atc tgt acc aac ctc    680
His His Glu Gln Pro Glu Thr Ser Asn Met Ser Ile Cys Thr Asn Leu
            200                 205                 210

tcc agc cgc tgg acc gtg ttc cag tcc agc atc ttc ggc gcc ttc gtg    728
Ser Ser Arg Trp Thr Val Phe Gln Ser Ser Ile Phe Gly Ala Phe Val
        215                 220                 225

gtc tac ctc gtg gtc ctg ctc tcc gta gcc ttc atg tgc tgg aac atg    776
Val Tyr Leu Val Val Leu Leu Ser Val Ala Phe Met Cys Trp Asn Met
230                 235                 240                 245

atg cag gtg ctc atg aaa agc cag aag ggc tcg ctg gcc ggg ggc acg    824
Met Gln Val Leu Met Lys Ser Gln Lys Gly Ser Leu Ala Gly Gly Thr
                250                 255                 260

cgg cct ccg cag ctg agg aag tcc gag agc gaa gag agc agg acc gcc    872
Arg Pro Pro Gln Leu Arg Lys Ser Glu Ser Glu Glu Ser Arg Thr Ala
                265                 270                 275

agg agg cag acc atc atc ttc ctg agg ctg att gtt gtg aca ttg gcc    920
Arg Arg Gln Thr Ile Ile Phe Leu Arg Leu Ile Val Val Thr Leu Ala
                280                 285                 290

gta tgc tgg atg ccc aac cag att cgg agg atc atg gct gcg gcc aaa    968
Val Cys Trp Met Pro Asn Gln Ile Arg Arg Ile Met Ala Ala Ala Lys
            295                 300                 305

ccc aag cac gac tgg acg agg tcc tac ttc cgg gcg tac atg atc ctc    1016
Pro Lys His Asp Trp Thr Arg Ser Tyr Phe Arg Ala Tyr Met Ile Leu
310                 315                 320                 325

ctc ccc ttc tcg gag acg ttt ttc tac ctc agc tcg gtc atc aac ccg    1064
Leu Pro Phe Ser Glu Thr Phe Phe Tyr Leu Ser Ser Val Ile Asn Pro
                330                 335                 340
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ctc ctg tac acg gtg tcc tcg cag cag ttt cgg cgg gtg ttc gtg cag      1112
Leu Leu Tyr Thr Val Ser Ser Gln Gln Phe Arg Arg Val Phe Val Gln
            345             350                 355

gtg ctg tgc tgc cgc ctg tcg ctg cag cac gcc aac cac gag aag cgc      1160
Val Leu Cys Cys Arg Leu Ser Leu Gln His Ala Asn His Glu Lys Arg
        360             365             370

ctg cgc gta cat gcg cac tcc acc acc gac agc gcc cgc ttt gtg cag      1208
Leu Arg Val His Ala His Ser Thr Thr Asp Ser Ala Arg Phe Val Gln
        375             380             385

cgc ccg ttg ctc ttc gcg tcc cgg cgc cag tcc tct gca agg aga act      1256
Arg Pro Leu Leu Phe Ala Ser Arg Arg Gln Ser Ser Ala Arg Arg Thr
390             395                 400                 405

gag aag att ttc tta agc act ttt cag agc gag gcc gag ccc cag tct      1304
Glu Lys Ile Phe Leu Ser Thr Phe Gln Ser Glu Ala Glu Pro Gln Ser
                410             415                 420

aag tcc cag tca ttg agt ctc gag tca cta gag ccc aac tca ggc gcg      1352
Lys Ser Gln Ser Leu Ser Leu Glu Ser Leu Glu Pro Asn Ser Gly Ala
                425             430                 435

aaa cca gcc aat tct gct gca gag aat ggt ttt cag gag cat gaa gtt      1400
Lys Pro Ala Asn Ser Ala Ala Glu Asn Gly Phe Gln Glu His Glu Val
        440             445             450

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